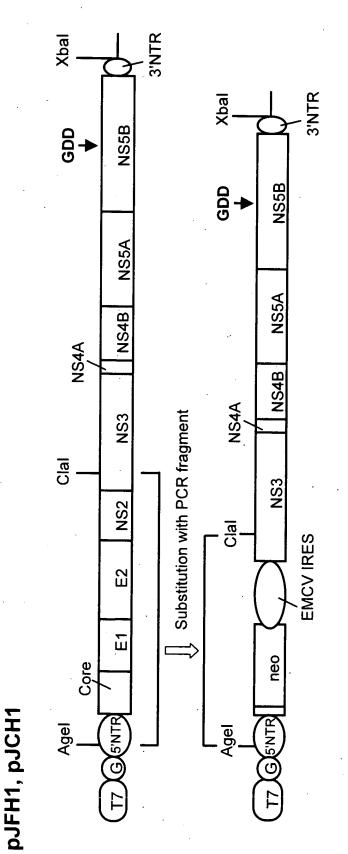
Fig.1



pSGREP-JFH1, pSGREP-JCH1

Fig.2A

10	20	30	40	50	60 GAACUACUGU
		90			
CUUCACGCAG	AAAGOGCCUA	GCCAUGGCGU	UAGUAUGAGU	GUCGUACAGC	CUCCAGGCCC
ccccccccccc	140	150	160	170	180
	GGAGAGCCAU	AGUGGUCUGC	GGAACCGGUG	AGUACACCGG	AAUUGCCGGG
190	200	AUAAACCCAC	220	230	240
AAGACUGGGU	CCUUUCUUGG		UCUAUGCCCG	GCCAUUUGGG	CGUGCCCCCG
250	260	270	260	290	300
CAAGACUGCU	AGOOGAGUAG	CGUUGGGUUG	CGAAAGGCCU	UGUGGUACUG	CCUGAUAGGG
310	320	330	340	350	360
CCCUUCCGAG	UGCCCCGGGA	GGUCUCGUAG	ACCGUGCACC	AUGAGCACAA	AUCCUAAACC
UCAAAGAAAA	ACCAAAAGAA	390 ACACCAACCG	UOGCCCAAUG	AUUGAACAAG	AUGGAUUGCA
430	440	450	460	470	480
CGCAGGUUCU	COGGCCGCUU	GGGUGGAGAG	GCUAUUCGGC	UAUGACUGGG	CACAACAGAC
490	500	510	520	530	540
AAUCGGCUGC	UCUGAUGOOG		GCUGUCAGOG	CAGGGGGGCC	COGUUCUUUU
550	560	570	580	590	600
UGUCAAGACC	GACCUGUCCG	GUGCCCUGAA	UGAACUGCAG	GACGAGGCAG	CGCGGCUAUC
GUGGCUGGCC	620	630	640	650	660
	AGGACGGGCG	UUCCUUGOGC	AGCUGUGCUC	GACGUUGUCA	CUGAAGOGGG
		690 GOGAAGUGOC			
730	740	750	760	770	780
UCCUGCOGAG	AAAGUAUCCA	UCAUGGCUGA	UGCAAUGCGG	COGCUGCAUA	CGCUUGAUCC
GGCUACCUGC	CCAUUCGACC	810 ACCAAGOGAA	ACAUCGCAUC	GAGCGAGCAC	GUACUCGGAU
		870 AGGAUGAUCU			
910	920	930	940	950	960
CGAACUGUUC	GOCAGGCUCA	AGGOGCGCAU	GCCCGACGGC	GAGGAUCUCG	UCGUGACCCA
9.70	080	990	1000	1010	1020
UGGCGAUGCC	ADOODUUDOU	AUAUCAUGGU	GGAAAAUGGC	CCCUUUUCUG	GAUUCAUOGA
		1050 CGGACCGCUA			
		1110 AAUGGGCUGA			
		1170 CCUUCUAUCG			
1210	1220	. 1230	1240	1250	1260
CCUCUCCCUC	CCCCCCCCU	AACGUUACUG	GCCGAAGCCG	CUUGGAAUAA	GGCCGGUGUG
		1790 UCCACCAUAU			1320 AGGGCCCGGA
		1350 ACGAGCAUUC			

Fig.2B

•					
1390 UGCAAGGUCU	1400 GUUGAAUGUC	1410 GUGAAGGAAG	1420 CAGUUCCUCU	1430 GGAAGCUUCU	1440 UGAAGACAAA
1450 CAAOGUCUGU	1460 AGOGACCCUU	1470 UGCAGGCAGC	1480 GGAACCCCCC	1490 ACCUGGOGAC	1500 AGGUGCCUCU
					1560 CAGUGCCACG
1570 UUGUGAGUUG	1580 GAUAGUUGUG	1590 GAAAGAGUCA	1600 AAUGGCUCUC	1610 CUCAAGCGUA	1620 UUCAACAAGG
1630 GGCUGAAGGA	1540 UGCCCAGAAG	1650 GUACOCCAUU	1660 GUAUGGGAUC	1670 UGAUCUGGGG	1680 CCUOGGUGCA
1690 CAUGCUUUAC	1700 AUGUGUUUAG	1710 UCGAGGUUAA	1720 AAAAAOGUCU	1730 AGGCCCCCCG	1740 AACCACGGGG
1750	1760	1770 ACACGAUGAU	1780	1790	1800
1810	1820	1830 CCCAUAGUG	1840	1850	1860
1870	1880	1890 CCUGUCCACA	1900	1910	1920
1930	1940	1950 UUACCACGGA	1960	1970	1960
1990	2000	2010 CUCGAGUGCU	2020	2030	2040
2050	2060	2070 GCCGUGCAAG	2080	2090	2100
2110	2120	2130 GGCUCGGAGA	2140	2150	2160
2170	2180	2190 GAAGGGGUCC	2200	2210	2220
2230	2240	2250 AGCAGCUGUG	2260	2276	2280
2290	2300	2310 CGACGUUGUU	2320	2330	2340
2350	2360	2370	2380	2390	2400
2410	2420	2430 UGUCGCGUAU	2440	2450	2460
2470	2480	2490 CCUGGGGUUU	2500	2510	2520
2530	2540	2550	.2560	2570	2580
2590	. 2600	AGUCAGGACC 2610	2620	2630	2640
ACAUAUGGCA	AAUUUCUOGC	OGAUGGGGGC	UGCGCUAGCG	GOGCCUAUGA	CAUCAUCAUA
		2670 GGAUGCUACC			
		2730 CAGACUAACU			

Fig.2C

2820 0000	UGAGAU	2810 GGOGGGAGGG	Z800 GUAGGCCUCG	2790 UAUAGAAGAG	2780 COCAUCOGA	2770 GUGACAACCC
2880 20GC	GAUUUU	2870 GGAGACACCU	2860 AUCAAGGGAG	convicorec 3820	2840 GGGGGAUUCC	2830 UUCUAUGGGA
3940 JGCC	CUUGAA	2930 GGGGCAUGGG	2920 GOGGCCCUUC	2910 CGAGCUCGCG	2900 AAAAGUGUGA	2890 CACUCAAAGA
3000 3GUC	UGUGGU	2990 CUCAGGGAGA	2980 AUAAUACCAG	2970 GGACGUCUCC	2960 AUAGAGGGUU	2950 GUGGCAUACU
3060	_	3050	3040	3030 GACGGGGUAC	3020	3010
812 0		3110	3100	3090 UGUCGACUUC	3080	3070
os r		3170	3160	3150 UGUCUCACGC	3140	3130
3240	:	3230	3220	3210 UGUUUCCACU	3200	3190
300	:	3290	3280	3270 CGACGCAGGG	3260	3250
360	1	3350	3340	3330 GUAUUUCAAC	3320	3310
420	· .	OTAT	3400	3390 UUUCACCGGC	3380	3370
480	3	3470	3460	3450 GAACUUCGCG	3440	3430
540	3	3530	3520	3510 CCCGUCCUGG	3500	3490
600	3	3590	3580	3570 CACACCUCUC	3550	3550
650 -	. 3	3650	3640	3630 GACGAAGUAC	3620	3610
720	3	3710	3700	3690 CCUAGCUGGA	3680	3670
780	3	3770	3760	3750 CAUCAUCGGC	3740	3730
840	3	3830	3820	3810 GUAUGAGGCU	3800	3790
ù à n		3890	3.880	3870 GCAGCGGAUA	3860	3850
960	3	. 3950	3940	3930 GCAGGCCCAG	3920	3910-
020	. 4	4010	4000	3990	3980	3970
	•			GGCCAGACAC 4050		
)80 \GU	GGCAUUÇ	4070 CUUCCAUGAU	CCCGCGGUGG	GCCAGGGAAC	UGUCAACACU	CUCGCAGGAU
				4110 GUCGACCAGU		

Fig.2D

4150 UGGUUAGCGU	4160 CCCAGAUCGC	4170 ACCACCGGG	4180 GGGCCACCG	4190 GCUUUGUCGU	4200 CAGUGGCCUG
4210 GUGGGGGCUG	4220 CCGUGGGCAG	4230 CAUAGGCCUG	4240 GGUAAGGUGC	4250 UGGUGGACAU	4250 CCUGGCAGGA
4270 UAUGGUGCGG	4280 GCAUUUCGGG	4290 GGCCUCGUC	4300 GCAUUCAAGA	4310 UCAUGUCUGG	4320 CGAGAAGCCC
4330 UCUAUGGAAG	4340 AUGUCAUCAA	4350 UCUACUGCCU	4360 GGGAUCCUGU	4370 CUCCGGGAGC	4380 CCUGGUGGUG
4390 GGGGUCAUCU	GCGCGGCCAU	4410 UCUGOGCOGC	4420 CACGUGGGAC	4430 CGGGGAGGG	4440 CGCGGUCCAA
4450 UGGAUGAACA	6460 GGCUUAUUGC	4470 CUUUGCUUCC	4480 AGAGGAAACC	4490 ACGUCGCCCC	4500 UACUCACUAC
4510 GUGACGGAGU	4520 CGGAUGCGUC	4530 GCAGCGUGUG	4540 ACCCAACUAC	4550 UUGGCUCUCU	4560 UACUAUAACC
AGCCUACUCA	GAAGACUCCA	4590 CAAUUGGAUA	ACUGAGGACU	GCCCCAUCCC	AUGCUCCGGA
UCCUGGCUCC	GCGACGUGUG	4650 GGACUGGGUU	UGCACCAUCU	UGACAGACUU	CAAAAAUUGG
CUGACCUCUA	AAUUGUUCCC	4710 CAAGCUGCCC	ecconcecen	UCAUCUCUUG	UCAAAAGGGG
UACANGGGUG	neneeccee	4770 CACUGGCAUC	AUGACCACGC	CCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	CGGCGCCAAC
AUCUCUGGCA	AUGUCCGCCU	4830 GGGCUCUAUG	AGGAUCACAG	GGCCUAAAAC	CUGCAUGAAC
ACCUGGCAGG	GGACCUUUCC	4890 UAUCAAUUGC	UACACGGAGG	GCCAGUGGGC	GCCGAAACCC
CCCACGNACU	ACAAGACCGC	4950 CAUCUGGAGG	GUGGCGCCU	OGGAGUAOGC	GGAGGUGACG
CAGCAUGGGU	OGUACUCCUA	5010 UGUAACAGGA	CUGACCACUG	ACAAUCUGAA	AAUUCCUUGC
CAACUACCUU	CUCCAGAGUU	5070 UUUCUCCUGG	GUGGACGGUG	UGCAGAUCCA	UAGGUUUGCA
5110 CCCACACCAA	5120 AGCOGUUUUU	5130 CCGGGAUGAG	5140 GUCUCGUUCU	5150 GCGUUGGGCU	5160 UAAUUCCUAU
GCUGUCGGGU	CCCAGCUUCC	5190 CUGUGAACCU	GAGCCCGACG	CAGACGUAUJI	GAGGUCCAUG
CUAACAGAUC	OGCCCCACAU	5250 CACGGCCGAG	ACUGOGGGGC -	GGCGCUUGGC	ACGGGGAUCA
5290 CCUCCAUCUG	5300 AGGCGAGCUC	5310 CUCAGUGAGC	5320 CAGCUAUCAG	5330 CACCGUCGCU	5340 GCGGGCCACC
5350 UGCACCACCC	5360 ACAGCAACAC	5370 CUAUGACGUG	5380 GACAUGGUCG	5390 AUGCCAACCU	5400 GCUCAUGGAG
5410 GGCGGUGUGG	5420 CUCAGACAGA	5430 GCCUGAGUCC	5440 AGGGUGCCCG	5450 UVCUGGACUU	5460 UCUCGAGCCA
5470 AUGGCCGAGG	5480 AAGAGAGCGA	5490 CCUUGAGCCC	5500 UCAAUACCAU	5510 CCGAGUGCAU	5520 GCUCCCCAGG

Fig.2E

5530	5540	5550	5560	SS70	5580
AGCGGGUUUC	CACGGGCCUU	ACCGGCUUGG	GCACGGCCUG	ACUACAACCC	GCCGCUCGUG
5590	5600	5610	5620	5630	5640
GAADOGUGGA	GGAGGCCAGA	UUACCAACOG	CCCACCGUUG	CUGGUUGUGC	DCUCCCCCC
5650	5660	5670	5680	5690	5700
CCCAAGAAGG	CCCCGACGCC	UCCCCCAAGG	AGACGCCGGA	CAGUGGGUCU	GÁGOGAGAGO
5710	5720	5730	5740	5750	5760
ACCAUAUCAG	AAGCCCUCCA	GCAACUGGCC	AUCAAGACCU	UUGGCCAGCC	CCCCUCGAGC
5770	5780	5790	5800	5810	5820
GGUGAUGCAG	GCUCGUCCAC	GGGGGGGGC	GCCGCCGAAU	CCGGCGGUCC	GAOGUCCOCU
5830	5840	5850	5860	5870	5880
GGUGAGCCGG	CCCCCCAGA	GACAGGUUCC	GCCUCCUCUA	UGCCCCCCU	CGAGGGGGAG
5890	5900	5910	5920	5930	5940
CCUGGAGAUC	CGGACCUGGA	GUCUGAUCAG	GUAGAGCUUC	AACCUCCCCC	CCAGGGGGG
5950	5960	5970	5980	5990	6000
GGGGUAGCUC	CCCCUUCGGG	CUCGGGGUCU	UGGUCUACUU	GCUCCGAGGA	GGACGAUACC
ACCEUGUGCU	GCUCCADGUC	AUACUCCUGG	ACCGGGGCUC	6050 UAAUAACUCC	CUGUAGOCCC
6070	6080	6090	6100	6110	6120
GAAGAGGAAA	AGUUGCCAAU	CAACCCUUUG	AGUAACUCGC	UGUUGOGAUA	CCAUAACAAG
				6170 AAAAGGUAAC	
ACGCAAGUGC	UCGACGCCCA	UUAUGACUCA	GUCUUAAAGG	6230 ACAUCAAGCU	AGCGGCUUCC
AAGGUCAGCG	CAAGGCUCCU	CACCUUGGAG	GAGGCGUGCC	AGUUGACUCC	
GCAAGAUCCA	AGUAUGGAUU	CGGGGCCAAG	GAGGUCCGCA	6350 GCUUGUCOGG	GAGGGCCGUU
AACCACAUCA	AGUCCGUGUG	GAAGGACCUC	CUGGAAGACC	6410 CACAAACACC	AAUUCCCACA
ACCAUCAUGG	CCAAAAAUGA	GGUGUUCUGC	GUGGACCCCG	6470 CCAAGGGGGG	UAAGAAACCA
6490	6500	6510	6520	6530	6540
GCUCGCCUCA	UCGUUUACCC	UGACCUCGGC	GUCCGGGUCU	GOGAGAAAAU	GGCCCUCUAU
GACAUUACAC	AAAAGCUUCC	UCAGGCGGUA	AUGGGAGCUU	6590 CCUAUGGCUU	CCAGUACUCC
6610	6620	6630	6640	6650	6660
CCUGCCCAAC	GGGUGGAGUA	UCÚCUUGAAA	GCAUGGGCGG	AAAAGAAGGA	CCCCAUGGGU
6670	6680	6690	6700	6710	6720
UUUUCGUAUG	AUACCCGAUG	CUUCGACUCA	ACCGUCACUG	AGAGAGACAU	CAGGACCGAG
				6770 GCACUGCCAU	
				6830 AGGGUCAAAC	
				6890 GUAACACCAU	6900 CACAUGCUAU

Fig.2F

6910 GUGAAAGCCC	6920 UAGCGGCCUG	6930 CAAGGCUGCG	6940 GGGAUAGUUG	6950 CCCCACAAU	
6970 GGCGAUGACC	6980 UAGUAGUCAU				
7030 AGAGCCUUCA	7040 CGGAGGCCAU	7050 GACCAGGUAC	7060 UCUGCCCCUC	7070 CUGGUGAUCC	
7090 GAAUAUGACC	7100 UGGAGCUAAU		7120 UCCUCAAAUG	7130 UGUCUGUGGC	7140 GUUGGGCCCG
7150 CGGGCCGCC	7160 GCAGAUACUA	7170 CCUGACCAGA	7180 GACCCAACCA	7190 CUCCACUCGC	
	7220 UUAGACACUC				
	7280 GGGUUCGCAU	•			
	7340 ACCAGAACCU				
7390 UUGGACCUUC 7450	7400 CAGCCAUAAU 7460	7410 UGAGAGGUUA 7470			
	ACGAACUGAC		7480 UCAGCCCUCA 7540	7490 GAAAACUUGG 7550	7500 GGCGCCACCC 7560
	GGAAGAGUCG 7580	GGCUCGCGCA	GUCAGGGCGU	CCCUCAUCUC	CCGUGGAGGG
AAAGCGGCCG	UUUGCGGCCG	7590 AUAUCUCUUC 7650	AAUUGGGCGG		
ACUCCAUTIGC	CGCAGGCGCG 7700	CCUACUGGAC	UUAUCCAGUU	GGUUCACOGU 7730	cccccccc
GGGGGGGACA	UUUUUCACAG	OGUGUOGOGC 7770	GCCCGACCCC	GCUCAUUACU 7790	CUUCGGCCUA
	UCGUAGGGGU	AGGCCUCUUC 7830	CUACUCCCCC	CUCCGUAGAG	COGCACACAC
	CCAUAGCUAA	CUGUUCCUUU	7900	บบบบบบบบบ	ບບບບບບບບບ 7920
	CUUUUUUUU	7950	UUUCUUCCCU	UCUCAUCUUA	UUCUACUUUC
UUUCUUGGUG	GCUCCAUCUU	AGCCCUAGUC	ACGGCUAGCU	GUGAAAGGUC	CGUGAGCCGC
AUGACUGCAG	0008 AGAGUGCCGU	AACUGGUCUC	UCUGCAGAUC	AUGU	UPUB

Fig.3A

ACCOGCCCCU	20 AAUAGGGGCG	ACACUCOGCC	40 AUGAAUCACU	50 CCCCUGUGAG	60 GAACUACUGU
70	08	90	100	110	120
CUUCAOGCAG	AUSUDODAAA	GCCAUGGCGU	UAGUAUGAGU	GUCGUACAGO	CUCCAGGCCC
130	140	150	160	170	180
cccccucccs	GGAGAGCCAU	AGUGGUCUGC	GGAACCGGUG	AGUACACOGG	AAUUGCOGGG
	cconncance	210	220	230	240
	500	AUAAACCCAC	UCUAUGCCCG	GCCAUUUGGG	OGUGCCCCCCG
250	260	270	280	290	300
CAAGACUGCU	AGCCGAGUAG	CGUUGGGUUG	CGAAAGGCCU	UGUGGUACUG	CCUGAUAGGG
310	320	330	340	350	360
UGCUUGCGAG	UGCCCCGGGA	GGUCUCGUAG	ACCGUGCACC	AUGAGCACAA	AUCCCAAACC
370	360	390	400	410	420
UCAAAGAAAA	ACCAAAAGAA	ACACUAACCG	UCGCCCAAUG	AUUGAACAAG	AUGGAUUGCA
430	440	450	460	470	480
CGCAGGUUCU	CCCCCCCUU	GGGUGGAGAG	GCUAUUCGGC	UAUGACUGGG	CACAACAGAC
490	500	510	520	530	540
AAUCGGCUGC	UCUGAUGCCG	CCGUGUUCCG	GCUGUCAGCG	CAGGGGGGCC	CGGUUCUUUU
550	560	570	580	590	600
UGUCAAGACC	GACCUGUCOG	GUGCCCUGAA	UGAACUGCAG	GACGAGGCAG	CGCGGCUAUC
610	620	630	640	650	660
GUGGCUGGCC	ACGACCGCCG	UUCCUUGCGC	AGCUGUGCUC	GACGUUGUCA	CUGAAGCGGG
670	660	690	700	710	720
AAGGGACUGG	CUGCUAUUGG	GCGAAGUGCC	GGGGCAGGAU	CUCCUGUCAU	CUCACCUUGC
730	740	750	760	770	780
UCCUGCOGAG	AAAGUAUCCA	UCAUGGCUGA	UGCAAUGOGG	CGGCUGCAUA	CGCUUGAUCC
790	800	810	820	830	840
GGCUACCUGC	CCAUUCGACC	ACCAAGCGAA	ACAUCGCAUC	GAGCGAGCAC	GUACUCGGAU
GGAAGCCCGGU	\$60 CUUGUCGAUC	AGGAUGAUCU	GGACGAAGAG	CAUCAGGGGC	UCGCGCCAGC
910	920	930	940	950	960
CGAACUGUUC	GCCAGGCUCA	AGGCGCCAU	GCCCGACGGC	GAGGAUCUCG	UOGUGACCCA
970	980	990	1000	1010	1020
UGGCGAUGCC	UGCUUGCCGA	AUAUCAUGGU	GGAAAAUGGC	COCUUUUCUG	GAUUCAUCGA
2030	1040	1050	1060	1070	1080
CUGUGGOOGG	CUGGGUGUGG	CGGACCGCUA	UCAGGACAUA	GOGUUGGCUA	CCOGUGAUAU
UGCUGAAGAG	1100 CUUGGCGGCG	AAUGGGCUGA	COGCUUCCUC	GUGCUUUACG	GUAUCGCCGC
1150	1160	1170	1180	1190	1200
UCCCGAUUCG	CAGCGCAUCG	CCUUCUAUCG	CCUUCUUGAC	GAGUUCUUCU	GAGUUUAAAC
1210	1220	1230	1240	1250	1260
CCUCUCCCUC	ccccccccu	AACGUUACUG	GCCGAAGCCG	CUUGGAAUAA	GGCCGGUGUG
	1280 UAUGUUAUJU				
1330	1340 UGUCUUCUUG	1350	1360	1370	1380

Fig.3B

. 1390	1400	1410	1420	1430	1440
UGCAAGGUCU	GUUGAAUGUC	GUGAAGGAAG	CAGUUCCUCU	GGAAGCUUCU	UGAAGACAAA
CAACGUCUGU	AGCGACCCUU	UGCAGGCAGC	GGAACCCCCC	ACCUGGOGAC	1500 AGGUGCCUCU
1510	1520	1530	1540	1550	1560
GCGGCCAAAA	GCCACGUGUA	UAAGAUACAC	CUGCAAAGGO	GGCACAACCC	CAGUGOCAOS
	GAUAGUUGUG	GAAAGAGUCA	AAUGGOUCUC	CUCAAGOGUA	1620 UUCAACAAGG
1630	1640	1650	1660	1670	CCUCGGUGCA
GGCUGAAGGA	UGCCCAGAAG	GUACCCCAUU	GUAUGGGAUC	UGAUCUGGGG	
CAUGCUUUAC	AUGUGUUUAG	UCGAGGUUAA	AAAAACGUCU	AGGCCCCCCCG	1740 AACCACGGGG
ACGUGGUUUU	CCUUUGAAAA	ACACGAUAAU	ACCAUGGCCC	CCAUCACCCC	
1810	1820	1830	1840	.1850	1860
CAGACACGAG	GUCUCUUGGG	CUCUAUAGUG	GUGAGCAUGA	CGGGGCGUGA	CAAGACAGAA
CAGGCCGGGG	AGGUCCAAGU	CCUGUCCACA	GUCACUCAGU	CCUUCCUCGG	1920 AACAUCCAUU
UCGGGGGUCU	1940	1950	1960	1970	1980
	UAUGGACUGU	UUACCACGGA	GCUGGCAACA	AGACACUAGC	CGGCUCGCGG
GGCCCGGUCA	CGCAGAUGUA	2010 CUCGAGCGCC	GAGGGGGACU	DGGUCGGGUG	GCCCAGCCCU
CCUGGGACCA	AAUCUUUGGA	2070 GCCGUGUACG	UGUGGAGCGG	UCGACCUGUA	UUUGGUCACG
2110	AUGUCAUCCC	2130	2140	2150	2160
CGGAACGCUG		GCCUCGAAGA	CGCGGGGACA	AGCGGGGAGC	GCUGCUCUCC
CCGAGACCCC	UUUCGACCUU	2190 GAAGGGGUCC	UCGGGGGGAC	CUGUGCUUUG	
2230	2240	2250	2260	2270	2280
CACGCUGUCG	GAAUCUUCOG	GGCAGCUGUG	UGCUCUCGGG	GUGUGGCUAA	GUCCAUAGAU
2290	2300	2310	ACGCGGUCUC	2330	2340
UUCAUCCCCG	UUGAGACGCU	OGACAUCGUC		CCACCUUUAG	UGACAACAGC
2350	2360	2370	2380	2390	2400
ACACCACCAG	CUGUGCCCCA	GACCUAUCAG	GUGGGGUACU	UGCACGCCCC	CACUGGCAGU
2410	2420	2430	2440	2450	CCUCCUCCUC
GGAAAAAGCA	CCAAGGUCCC	OGUCGOGUAC	GCCGCCAGG	GGUAUAAAGU	
2470	2480	2490	2500	2510	2520
AAUCCCUCGG	UGGCUGCCAC	CCUGGGAUUU	GGGGCGUACU	UGUCCAAGGC	ACAUGGCAUC
2530	2540	2550	2560	2570	2580
AACCCCAACA	UUAGGACUGG	AGUCAGAACU	GUGACGACCG	GGGAGCOCAU	UACAUACUCC
2590	2600	2610	2620	2630	2640
ACGUAUGGUA	AAUUCCUCGC	CGAUGGGGGC	UGCGCAGGCG	GOGCCUAUGA	CAUCAUCAUA
2650	2660	2670	2680	2690	2700
UGOGAUGAAU	GCCACUCUGU	GGAUGCUACC	ACUAUUCUCG	GCAUCGGAC	AGUCCURGAC
. 2710	2720	2730 CAGGCUAACU	2740	2750	2760

Fig.3C

2770 GUGACAACOC	2780 COCAUCOCAA	2790 UAUAGAGGAG	2800 GUAGCCCUCG	2810 GACAGGAGGG	2820 UGAGAUCCCC
2830 UUCUAUGGGA	2840 GGGCGUUUCC	2850 CCUGUCUUAC	2850 AUCAAGGGAG	2870 GGAGGCACUU	2880 GAUUUUCUGC
2890 CACUCAAAGA	2900 AAAAGUGUGA	2910 CGAGCUCGCA	2920 ACGCCCCUC	2930 GGGGCAUGGG	2940 CUUGAACGCU
GUGGCAUAUU	ACAGAGGGUU	GGACGUCUCC	AUAAUACCAA	CUCAAGGAGA	3000 UGUGGUGGUC
GUUGOCACCG	ACGCCCUCAU	GACGGGGUAU	ACUGGAGACU		GAUCGACUGC
3070 AAOGUAGOGG	3080 UCACCCAGGC	3090 OGUAGACUUC	AGCCUGGACC	3110 CCACCUUCAC	3120 UAUAACCACA
CAGACUGUCC	CGCAAGACGC	UGUCUCAGGU	AGUCAGOGCC	GAGGGGGCAC	
AGACUGGGCA	UUUAUAGGUA	UGUUUCCACU	GGUGAGOGAG		GUUUGACAGU
GUAGUACUCU	GUGAGUGCUA	CGACGCAGGA	ecnecnneen	3290 AUGAGCUCUC	ACCAGUGGAG
ACGACCGUCA	GCCUCAGGGC	GUAUUUCAAC	ACGCCUGGCU	UGCCUGUGUG	CCAGGACCAC
CUUGAGUUUU	GGGAGGCAGU	UUUCACCGGC	CUCACACACA	3410 UAGACGCUCA	UUUCCUUUCC
CAGACAAAGC	AGUCGGGGGA	AAAUUUOGCA	UACUUAGUAG	3470 CCUAUCAGGC	CACAGUGUGC
GCCAGGGCCA	AAGCGCCCCC	CCCGUCCUGG	GACGUCAUGU	3530 GGAAGUGCUU	GACUCGACUC
AAGCCCACGC	UUGUGGGCCC	UACACCUCUC	CUGUACOGUU	3590 UGGGCUCUGU	UACCAACGAG
GUCACOCUUA	CACACCOOGU	GACAAAAUAC	AUCGCCACAU	3650 GCAUGCAAGC	UGACCUCGAG
GUCAUGACCA	GCACGUGGGU	CCUGGCUGGG	GGAGUCUUAG	3710 CAGCCGUCGC	CGCGUAUUGC
UUAGOGACOG	GGUGUGUUUC	CAUCAUUGGC	CGUUUACACA	3770 UCAACCAGCG	AGCUGUCGUC
GCUCOGGACA	AGGAGGUCCU	CUAUGAGGCU	UUUGAUGAGA	3830 UGGAGGAAUG	UGCCUCCAGA
ececcncncc	UUGAAGAGGG	GCAGCGGAUA	GCCGAGAUGC	3890 UGAAGUCCAA	GAUCCAAGGC
UUAUUGCAGC	AAGCCUCUAA	ACAGGCCCAG	GACAUACAAC		AGCUUCGUGG
3970 CCCAAGAUGG	3980 AGCAAUUCUG	3990 GGCCAAACAU	4000 AUGUGGAACU	4010 UCAUAAGOGG	4020 CAUUCAGUAC
				4070 CUUCCAUGAU	
				4130 UUCUUAACAU	4140 UCUGGGGGC

Fig.3D

4200	4190	4180	4170	4160	4150
CAGUGGCCUG	GCUUUGUUGU	GGGCCACUG	GCCACCOGCG	CCCAAAUUGC	UGGCUGGCGU
4260	4250	4240	4230	4220	4210
CCUGGCAGGG	UGGUGGACAU	GGUAAAGUGC	CAUAGGCUUG	CUGUUGGCAG	GUGGGAGCUG
4320 CGAGAAGCCC					
4380	4370	4360	4350	4340	4330
UCUGGUGGUG	CUCCAGGUGC	GGGAUUCUGU	CUUGCUGCCU	AUGUCAUCAA	UCCAUGGAGG
4440	4430	4420	4410	4400	4390
CGCGGUCCAA	CGGGGGAAGG	CAUGUGGGAC	UCUGCGCCGC	GCGCGGCCAU	GGAGUCAUCU
4500 UACUCACUAC					
4560 CACUAUAACU					
4620	4610	4600	4590	4580	4570
AUGCGCCGGC	GCCCCAUCCC	ACUGAGGAUU	CAACUGGAUC	GGAGACUUCA	AGUCUACUCA
4580	4570	4660	4650	4640	4630
UAAGAACUGG	UAACAGACUU	UGUACCAUCC	GGACUGGGUC	GOGAUGUGUG	UCGUGGCUCC
4740	4730	4720	4710	4700	4690
CCAAAAGGGG	UUAUCUCUUG	GCCUCCCCU	AAAGAUGCCU	AGCUGUUCCC	CUGACCUCCA
4800	4790	4780	4770	4760	4750
CGGCGCCAAC	GAUGCCCCUG	AUGACCACAC	CACUGGCAUC	UGUGGGCCGG	UACAAGGGCG
4860	4850	4840	4830	4820	4810
CUGCAUGAAC	GACCCAAAAC	AGAAUCACAG	GGGCUCUAUG	ACGUCCGCUU	AUCUCUGGCA
4920	4910	4900	4890	4880	4870
GCCGAAACCC	GCCAGUGCUU	UAUACAGAAG	UAUCAAUUGU	GACCUUUCC	ACCUGGCAGG
4980	. 4970	4960	4950	4940	4930
GGAAGUGACG	CAGAGUACGC	GUGGCGGCCU	CAUCUGGAGA	UCAAGACCGC	GCGUUAAACU
5040	5030	5020	5010	5000	4990
AGUCCCUUGC	ACAACUUAAA	CUGACCACUG	UAUAACAGGG	CAUAUGCCUA	CAGCACGGAU
9100	5090	5080	5070	5060	5050
UAGGUCCGCC	UACAAAUOCA	GUGGACOGAG	UUUCUCUUGG	CUCCAGAGUU	CAACUCCCCU
5150	5150	5140	5130	5120	5110
CAAUUCAUUU	GCGUUGGGCU	GUCUCGUUCA	CCGGGAUGAG	AGCCGUUUUU	CCCACACCAA
5220	5210	5200	5190	5180	5170
GAUGUCCAUG	CUGAGGUAGU	GAGCCCGACA	CUGUGACCCU	CUCAGCUUCC	GUCGUCGGGU
5280 GCGGGGGUCA			5250 CACGGCGGAG		
5340 GCGAGCCACC					
			5370 CUAUGAUGUG		
5460 CCUCGACUCA			5430 GUCUGAGUCC		
5520 GCUCCCCAGG	5510	5500	5490	5480	5470
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Fig.3E

5530	5540	5550	5560	5570	5580
AAGAGGUUCC	CACCGGCCUU	ACCGGCUUGG	GOGGGCCUG	AUUACAACCC	ACCGCUUGUG
		5610 UUACCAACCA			
5650	5660	5670	5680	5690	GAGCGAGAGC
CCCAAAAAGA	CCCCGACGCC	UCCUCCAAGG	AGACGCCGGA	CAGUGGGUCU	
ACCAUAGGAG	AUGCCCUCCA	5730 ACAGCUGGCC	AUCAAGUCCU	UUGGCCAGCC	CCCCCCAAGC
5770	5760	5790	GCCGCCGACU	5810	5820
GGCGAUUCAG	GCCUTUCCAC	GGGGGGGAC		COGGOGAUCG	GACACCCCCU
5830	5840	5850	5860	UGCCCCCCCU	5880
GACGAGUUGG	CUCUUUCGGA	GACAGGUUCU	ACCUCCUCCA		CGAGGGGGAG
CCUGGGGACC	CAGACCUGGA	5910 GCCUGAGCAG	GUAGAGCUUC	AACCUCCUCC	CCAGGGGGGG
5950	5960	5970	5980	5990	6000
GAGGCAGCUC	COGGCUCGGA	CUCGGGGUCC	UGGUCUACUU	GCUCCGAGGA	GGAUGACUCC
6010	6020	6030	6040	6050	060
GUCGUGUGCU	GCUCCAUGUC	AUAUUCCUGG	ACCGGGGCUC	UAAUAACUCC	UUGUAGCCCC
		6090 UAACUCCUUG			
GUAUACUGUA	6140	6150	6160	6170	6180
	CUACAUCAAA	GAGUGCCUCA	CUAAGGGCUA	AAAAGGUAAC	UUUUGAUAGG
6190	6200	6210	6220	5230	6240
AUGCAAGUGC	UOGACGCCUA	UUAUGAUUCA	GUCUUAAAGG	ACAUCAAGCU	AGCGGCCUCC
5250	6260	6270	6280	6290	ACCCCACUCU
AAGGUCAGCG	CAAGGCUCCU	CACCUUAGAG	GAGGCGUGCC	AAUUGACCCC	
6310 GCAAGAUCCA	6320 AGUAUGGGUU	6330 UGGGGCUAAG	6340 GAGGUCCGCA		
6370	6380	6390	6400	6410	6420
AACCACAUCA	AGUCCGUGUG	GAAGGACCUC	UUGGAAGACU	CACAAACACC	AAUUCCUACA
6430	6440	6450	6460	6470	6480
ACCAUCAUGG	CCAAAAAUGA	GGUGUUCUGC	GUGGACCCCG	CCAAGGGGGG	UAAAAAACCA
6490	6500	6510	6520	6530	6540
GCUCGCCUUA	UCGUUUACCC	UGACCUCGGC	GUCAGGGUCU	Gogagaagau	GGCCCUUUAU
6550	6560	6570	6580	6590	6600
GAUGUCACAC	AAAAGCUUCC	UCAGGCGGUG	AUGGGGGCUU	CUUAUGGCUU	CCAGUACUCC
		6530 UCUCUUGAAG			
		6690 CUUUGACUCA			
		6750 CUCCUUACCC			
		6810 AGGGCCCAUG			
		6870 GGUGCUUACC			

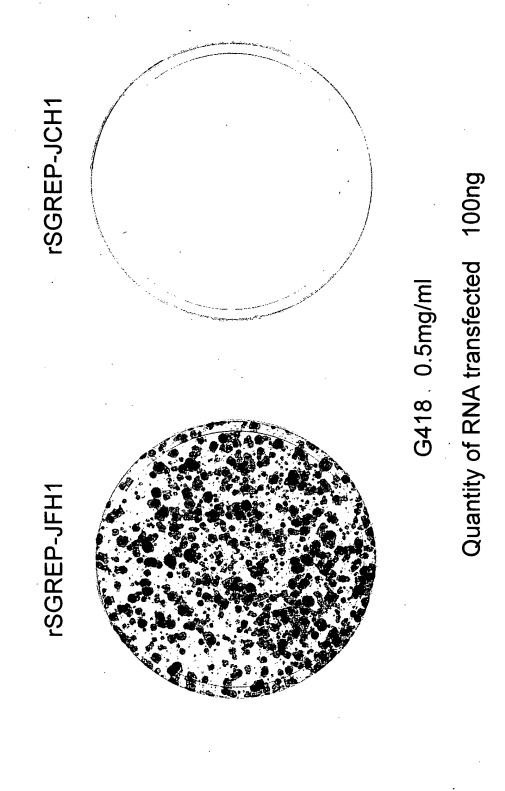
Fig.3F

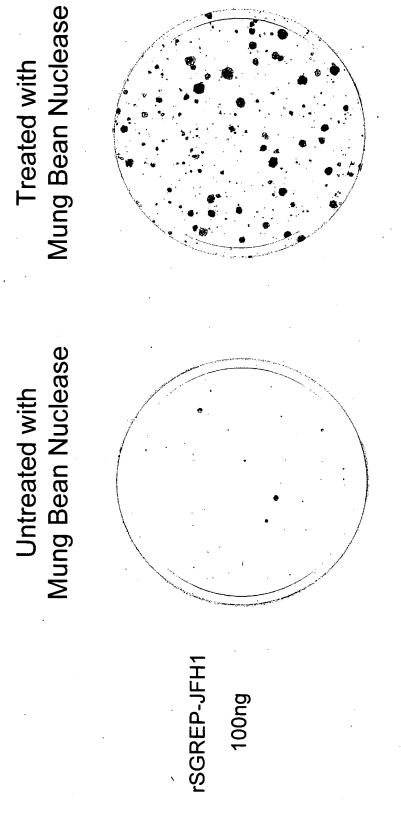
6910 GUAAAAGCCC	6920 UAGCGGCUUG	6930 CAAGGCUGCG		OGCCCACGAU	GCUGGUAUGC
6970 GGOGACGACU	6980 UGGUCGUCAU	6990 CUCAGAAAGC	7000 CAGGGGACUG	7010 AGGAGGACGA	7020 GOGGAACCUG
7030 AGAGCCUUCA	7040 CGGAGGCUAU	7050 GACCAGGUAU	7060 UCUGCCCCUC	7070 CUGGUGACCC	7080 CCCCAGACOG
7090 GAAUAUGACC	7100 UGGAGCUAAU		7120 UCCUCAAACG		7140 ACUUGGCCCA
7150 CAGGCCCCCC	7160 GCAGAUACUA	7170 CCUGACCAGA	7180 GACCCCACCA	7190 CUUCAAUUGC	7200 CCGGGCUGCC
7210 UGGGAAACAG	7220 UUAGACACUC				•
7270 CCAACCAUAU	7280 GGGUUCGCAU	GGUCCUGAUG		7310 UCUCCAUUCU	
7330 GACACCCUAG	7340 ACCAGAACCU	UAACUUUGAA	AUGUACGGAU	7370 COGUGUACUC	
7390 CUGGACCUCC	7400 CAGCCAUAAU	UGAAAGGUUA	CACGGGCUUG	ACGCCUUCUC	
7450 UACACUCCCC	ACGAACUGAC	GCGGGUGGCU		GAAAACUTUGG	
•	GGAAGAGUCG		GUUAGGGCGU	CCCACACAC	
AGGGCGGCCG	7580 UUUGOGGUCG	GUACCUCUUC	AACUGGGCGG	UGAAGACCAA	GCUCAAACUC
ACUCCUUUGC	7640 CGGAGGCACG	CCUCCUGGAU	UUGUCCAGUU	GGUUUACCGU	COCCCCCCCCC
GGGGGCGACA	7700 UUUAUCACAG	CCUCUCGCGu	GCCCGACCCC	GCCUAUUACU	CCUUAGCCUA
	7760 cuguageggu				
UAGCUACACU	7820 CCAUAGCUAA	COCCOCCO	OOOOOOOO	บบบบบบบบบบบ	บบบบบบบบบบ
	CUUUUUUUU	UUUUUCCCUC		UCUCAUCUUA	UUCUACUUUC
UUUCUUGGUG	7940 GCUCCAUCUU	AGCCCUAGUC	ACGGCUAGCU	GUGAAAGGUC	CGUGAGCCGC
7990 AUGACUGCAG	8000 AGAGUGCCGU	8010 AACUGGUCUC	8020 UCUGCAGAUC	AUGU	8040

rSGREP -JFH1/GND rSGREP -JFH1/dGDD rSGREP-JFH1 Quantity of RNA transfected 100ng 300ng Fig.4

14/25

G418 1.0mg/ml





G418 1.0mg/ml

